## SEQUENCE LISTING

<110>	Altmann, Scott W	
	Wang, Luquan	
	Graziano, Michael	
	Murgolo, Nick	
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ttc g Phe G																1248.
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agc ( Ser I								-								1488
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Val Thr Gly Glu His Leu Ala Leu Leu Gln Arg Ile Cys Pro Arg Leu 65 70 75 80

Tyr Asn Gly Pro Asn Thr Thr Phe Ala Cys Cys Ser Thr Lys Gln Leu 85 90 95

Leu Ser Leu Glu Ser Ser Met Ser Ile Thr Lys Ala Leu Leu Thr Arg Cys Pro Ala Cys Ser Asp Asn Phe Val Ser Leu His Cys His Asn Thr Cys Ser Pro Asp Gln Ser Leu Phe Ile Asn Val Thr Arg Val Val Glu Arg Gly Ala Gly Glu Pro Pro Ala Val Val Ala Tyr Glu Ala Phe Tyr Gln Arg Ser Phe Ala Glu Lys Ala Tyr Glu Ser Cys Ser Gln Val Arg Ile Pro Ala Ala Ala Ser Leu Ala Val Gly Ser Met Cys Gly Val Tyr Gly Ser Ala Leu Cys Asn Ala Gln Arg Trp Leu Asn Phe Gln Gly Asp Thr Gly Asn Gly Leu Ala Pro Leu Asp Ile Thr Phe His Leu Leu Glu Pro Gly Gln Ala Leu Pro Asp Gly Ile Gln Pro Leu Asn Gly Lys Ile Ala Pro Cys Asn Glu Ser Gln Gly Asp Asp Ser Ala Val Cys Ser Cys Gln Asp Cys Ala Ala Ser Cys Pro Val Ile Pro Pro Pro Glu Ala Leu Arg Pro Ser Phe Tyr Met Gly Arg Met Pro Gly Trp Leu Ala Leu Ile Ile Ile Phe Thr Ala Val Phe Val Leu Leu Ser Ala Val Leu Val Arq Leu Arg Val Val Ser Asn Arg Asn Lys Asn Lys Ala Glu Gly Pro Gln Glu Ala Pro Lys Leu Pro His Lys His Lys Leu Ser Pro His Thr Ile 

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Gly	Leu 370	Thr	Phe	Ile	Glu	Leu 375	Thr	Thr	Asp	Pro	Val 380	Glu	Leu	Trp	Ser
Ala 385	Pro	Lys	Ser	Gln	Ala 390	Arg	Lys	Glu	Lys	Ser 395	Phe	His	Asp	Glu	His 400
Phe	Gly	Pro	Phe	Phe 405	Arg	Thr	Asn	Gln	Ile 410	Phe	Val	Thr	Ala	Arg 415	Asn
Arg	Ser	Ser	Tyr 420	Lys	Tyr	Asp	Ser	Leu 425	Leu	Leu	Gly	Ser	Lys 430	Asn	Phe
Ser	Gly	Ile 435	Leu	Ser	Leu	Asp	Phe 440	Leu	Leu	Glu	Leu	Leu 445	Glu	Leu	Gln
Glu	Arg 450	Leu	Arg	His	Leu	Gln 455	Val	Trp	Ser	Pro	Glu 460	Ala	Glu	Arg	Asn
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Thr	Ser	Leu 515	Val	Asp	Trp	Lys	Asp 520	His	Phe	Leu	Tyr	Cys 525	Ala	Asn	Ala
Pro	Leu 530	Thr	Phe	Lys	Asp	Gly 535	Thr	Ser	Leu	Ala	Leu 540	Ser	Cys	Met	Ala
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580 585

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- Cys Asn Ile Asn Val Gly Leu Asp Gln Glu Leu Ala Leu Pro Lys Asp 865 870 875 880
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Gln Tyr Pro Ser Pro Ala Asp Ala Asp Ala Asn Val Asn Tyr Gly 1295 1300 1305 Phe Ala Pro Glu Leu Ala His Gly Ala Asn Ala Ala Arg Ser Ser 1310 1315 1320 Leu Pro Lys Ser Asp Gln Lys Phe <210> 3 <211> 3999 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (1)..(3999) <223> <400> 3 atg gcg gag gcc ggc ctg agg ggc tgg ctg ctg tgg gcc ctg ctc ctg 48 Met Ala Glu Ala Gly Leu Arg Gly Trp Leu Leu Trp Ala Leu Leu Leu 10 ege ttg gee cag aqt gag eet tae aca ace ate cae caq eet qqe tae 96 Arg Leu Ala Gln Ser Glu Pro Tyr Thr Thr Ile His Gln Pro Gly Tyr tge gee tte tat gae gaa tgt ggg aag aac cea gag etg tet gga age 144 Cys Ala Phe Tyr Asp Glu Cys Gly Lys Asn Pro Glu Leu Ser Gly Ser 35 40 ctc atg aca ctc tcc aac gtg tcc tgc ctg tcc aac acg ccq qcc cqc 192 Leu Met Thr Leu Ser Asn Val Ser Cys Leu Ser Asn Thr Pro Ala Arg 50 55 aag atc aca ggt gat cac ctg atc cta tta cag aag atc tgc ccc cqc 240 Lys Ile Thr Gly Asp His Leu Ile Leu Leu Gln Lys Ile Cys Pro Arg 65 70 ctc tac acc ggc ccc aac acc caa gcc tgc tgc tcc gcc aag cag ctg 288 Leu Tyr Thr Gly Pro Asn Thr Gln Ala Cys Cys Ser Ala Lys Gln Leu 85 gta tca ctg gaa gcg agt ctg tcg atc acc aag gcc ctc ctc acc cgc 336 Val Ser Leu Glu Ala Ser Leu Ser Ile Thr Lys Ala Leu Leu Thr Arg

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		_	_	_		Cys		-				_	_	aat Asn		1440
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tgg gac Trp Asp 116	Ile S	_		_	_					_		_	3519
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Leu Met Thr Leu Ser Asn Val Ser Cys Leu Ser Asn Thr Pro Ala Arg 50 55 60

Lys Ile Thr Gly Asp His Leu Ile Leu Leu Gln Lys Ile Cys Pro Arg 65 70 75 80

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Val Ser Leu Glu Ala Ser Leu Ser Ile Thr Lys Ala Leu Leu Thr Arg 100 105 110

Cys Pro Ala Cys Ser Asp Asn Phe Val Asn Leu His Cys His Asn Thr 115 120 125

Cys Ser Pro Asn Gln Ser Leu Phe Ile Asn Val Thr Arg Val Ala Gln 130 135 140

Gln His Ser Phe Ala Glu Gln Ser Tyr Asp Ser Cys Ser Arg Val Arg

165 170 175

Val Pro Ala Ala Ala Thr Leu Ala Val Gly Thr Met Cys Gly Val Tyr 180 185 190

Gly Ser Ala Leu Cys Asn Ala Gln Arg Trp Leu Asn Phe Gln Gly Asp

Thr Gly Asn Gly Leu Ala Pro Leu Asp Ile Thr Phe His Leu Leu Glu 210 215 220

Pro Gly Gln Ala Val Gly Ser Gly Ile Gln Pro Leu Asn Glu Gly Val 225 230 235

Ala Arg Cys Asn Glu Ser Gln Gly Asp Asp Val Ala Thr Cys Ser Cys 245 250 255

Gln Asp Cys Ala Ala Ser Cys Pro Ala Ile Ala Arg Pro Gln Ala Leu 260 265 270

Asp Ser Thr Phe Tyr Leu Gly Gln Met Pro Gly Ser Leu Val Leu Ile 275 280 285

Ile Ile Leu Cys Ser Val Phe Ala Val Val Thr Ile Leu Leu Val Gly 290 295 300

Phe Arg Val Ala Pro Ala Arg Asp Lys Ser Lys Met Val Asp Pro Lys 305 310 315 320

Lys Gly Thr Ser Leu Ser Asp Lys Leu Ser Phe Ser Thr His Thr Leu 325 330 335

Leu Gly Gln Phe Phe Gln Gly Trp Gly Thr Trp Val Ala Ser Trp Pro 340 345 350

Leu Thr Ile Leu Val Leu Ser Val Ile Pro Val Val Ala Leu Ala Ala 355 360 365

Gly Leu Val Phe Thr Glu Leu Thr Thr Asp Pro Val Glu Leu Trp Ser 370 375 380

Ala Pro Asn Ser Gln Ala Arg Ser Glu Lys Ala Phe His Asp Gln His 385 390 395 400

Phe Gly Pro Phe Phe Arg Thr Asn Gln Val Ile Leu Thr Ala Pro Asn 405 410 415

Arg Ser Ser Tyr Arg Tyr Asp Ser Leu Leu Gly Pro Lys Asn Phe Ser Gly Ile Leu Asp Leu Asp Leu Leu Glu Leu Glu Leu Gln Glu Arg Leu Arg His Leu Gln Val Trp Ser Pro Glu Ala Gln Arg Asn Ile Ser Leu Gln Asp Ile Cys Tyr Ala Pro Leu Asn Pro Asp Asn Thr Ser Leu Tyr Asp Cys Cys Ile Asn Ser Leu Leu Gln Tyr Phe Gln Asn Asn Arg Thr Leu Leu Leu Thr Ala Asn Gln Thr Leu Met Gly Gln Thr Ser Gln Val Asp Trp Lys Asp His Phe Leu Tyr Cys Ala Asn Ala Pro Leu Thr Phe Lys Asp Gly Thr Ala Leu Ala Leu Ser Cys Met Ala Asp Tyr Gly Ala Pro Val Phe Pro Phe Leu Ala Ile Gly Gly Tyr Lys Gly Lys Asp Tyr Ser Glu Ala Glu Ala Leu Ile Met Thr Phe Ser Leu Asn Asn Tyr Pro Ala Gly Asp Pro Arg Leu Ala Gln Ala Lys Leu Trp Glu Glu Ala Phe Leu Glu Glu Met Arg Ala Phe Gln Arg Arg Met Ala Gly Met Phe Gln Val Thr Phe Thr Ala Glu Arg Ser Leu Glu Asp Glu Ile Asn Arg Thr Thr Ala Glu Asp Leu Pro Ile Phe Ala Thr Ser Tyr 

Ile Val Ile Phe Leu Tyr Ile Ser Leu Ala Leu Gly Ser Tyr Ser Ser

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Leu 705	Val	Leu	Ser	Val	Gly 710	Ala	Asp	Asn	Ile	Phe 715	Ile	Phe	Val	Leu	Glu 720
Tyr	Gln	Arg	Leu	Pro 725	Arg	Arg	Pro	Gly	Glu 730	Pro	Arg	Glu	Val	His 735	Ile
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Val	Arg 770	Thr	Phe	Ala	Leu	Thr 775	Ser	Gly	Leu	Ala	Val 780	Ile	Leu	Asp	Phe
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Cys 865	His	Ile	Ser	Val	Gly 870	Leu	Asp	Gln	Glu	Leu 875	Ala	Ļeu	Pro	Lys	Asp 880
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- Glu Ala Gly Met Asn Ala Ile Cys Ser Ser Ala Gly Cys Asn Asn Phe 915 920 925
- Ser Phe Thr Gln Lys Ile Gln Tyr Ala Thr Glu Phe Pro Glu Gln Ser 930 935 940
- Tyr Leu Ala Ile Pro Ala Ser Ser Trp Val Asp Asp Phe Ile Asp Trp 945 950 955 960
- Leu Thr Pro Ser Ser Cys Cys Arg Leu Tyr Ile Ser Gly Pro Asn Lys 965 970 975
- Asp Lys Phe Cys Pro Ser Thr Val Asn Ser Leu Asn Cys Leu Lys Asn 980 985 990
- Cys Met Ser Ile Thr Met Gly Ser Val Arg Pro Ser Val Glu Gln Phe 995 1000 1005
- His Lys Tyr Leu Pro Trp Phe Leu Asn Asp Arg Pro Asn Ile Lys 1010 1015 1020
- Cys Pro Lys Gly Gly Leu Ala Ala Tyr Ser Thr Ser Val Asn Leu 1025 1030 1035
- Thr Ser Asp Gly Gln Val Leu Ala Ser Arg Phe Met Ala Tyr His 1040 1045 1050
- Lys Pro Leu Lys Asn Ser Gln Asp Tyr Thr Glu Ala Leu Arg Ala 1055 1060 1065
- Ala Arg Glu Leu Ala Ala Asn Ile Thr Ala Asp Leu Arg Lys Val 1070 1080
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- Asn Val Phe Tyr Glu Gln Tyr Leu Thr Ile Leu Pro Glu Gly Leu 1100 1105 1110
- Phe Met Leu Ser Leu Cys Leu Val Pro Thr Phe Ala Val Ser Cys 1115 1120 1125
- Leu Leu Leu Gly Leu Asp Leu Arg Ser Gly Leu Leu Asn Leu Leu

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Ser Ile Val Met Ile Leu Val Asp Thr Val Gly Phe Met Ala Leu 1145 1150 1155

Trp Asp Ile Ser Tyr Asn Ala Val Ser Leu Ile Asn Leu Val Ser 1160 1165 1170

Ala Val Gly Met Ser Val Glu Phe Val Ser His Ile Thr Arg Ser 1175 1180 1185

Phe Ala Ile Ser Thr Lys Pro Thr Trp Leu Glu Arg Ala Lys Glu 1190 1195 1200

Ala Thr Ile Ser Met Gly Ser Ala Val Phe Ala Gly Val Ala Met 1205 1210 1215

Thr Asn Leu Pro Gly Ile Leu Val Leu Gly Leu Ala Lys Ala Gln 1220 1225 1230

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Leu Gly Leu Leu His Gly Leu Val Phe Leu Pro Val Ile Leu Ser 1250 1255 1260

Tyr Val Gly Pro Asp Val Asn Pro Ala Leu Ala Leu Glu Gln Lys 1265 1270 1275

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<213> Mus sp.

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Ala S						gçc Ala 1075									3249
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Pro Leu Thr Tyr Lys Asp Gly Thr Ala Leu Ala Leu Ser Cys Ile Ala Asp Tyr Gly Ala Pro Val Phe Pro Phe Leu Ala Val Gly Gly Tyr Gln Gly Thr Asp Tyr Ser Glu Ala Glu Ala Leu Ile Ile Thr Phe Ser Ile Asn Asn Tyr Pro Ala Asp Asp Pro Arg Met Ala His Ala Lys Leu Trp Glu Glu Ala Phe Leu Lys Glu Met Gln Ser Phe Gln Arg Ser Thr Ala Asp Lys Phe Gln Ile Ala Phe Ser Ala Glu Arg Ser Leu Glu Asp Glu Ile Asn Arg Thr Thr Ile Gln Asp Leu Pro Val Phe Ala Ile Ser Tyr Leu Ile Val Phe Leu Tyr Ile Ser Leu Ala Leu Gly Ser Tyr Ser Arg Trp Ser Arg Val Ala Val Asp Ser Lys Ala Thr Leu Gly Leu Gly Gly Val Ala Val Val Leu Gly Ala Val Val Ala Ala Met Gly Phe Tyr Ser Tyr Leu Gly Val Pro Ser Ser Leu Val Ile Ile Gln Val Val Pro Phe Leu Val Leu Ala Val Gly Ala Asp Asn Ile Phe Ile Phe Val Leu Glu Tyr Gln Arg Leu Pro Arg Met Pro Gly Glu Gln Arg Glu Ala His Ile Gly Arg Thr Leu Gly Ser Val Ala Pro Ser Met Leu Leu Cys Ser Leu Ser Glu Ala Ile Cys Phe Phe Leu Gly Ala Leu Thr Ser Met Pro Ala

Val Arg Thr Phe Ala Leu Thr Ser Gly Leu Ala Ile Ile Phe Asp Phe
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Asn Leu Pro Pro Pro Lys Gln Lys Glu Gly Leu Leu Cys Phe Phe 820 825 830

Arg Lys Ile Tyr Thr Pro Phe Leu Leu His Arg Phe Ile Arg Pro Val 835 840 845

Val Leu Leu Leu Phe Leu Val Leu Phe Gly Ala Asn Leu Tyr Leu Met 850 855 860

Cys Asn Ile Ser Val Gly Leu Asp Gln Asp Leu Ala Leu Pro Lys Asp 865 870 875 880

Ser Tyr Leu Ile Asp Tyr Phe Leu Phe Leu Asn Arg Tyr Leu Glu Val 885 890 895

Gly Pro Pro Val Tyr Phe Asp Thr Thr Ser Gly Tyr Asn Phe Ser Thr 900 905 910

Glu Ala Gly Met Asn Ala Ile Cys Ser Ser Ala Gly Cys Glu Ser Phe 915 920 925

Ser Leu Thr Gln Lys Ile Gln Tyr Ala Ser Glu Phe Pro Asn Gln Ser 930 935 940

Tyr Val Ala Ile Ala Ala Ser Ser Trp Val Asp Asp Phe Ile Asp Trp 945 950 955 960

Leu Thr Pro Ser Ser Ser Cys Cys Arg Ile Tyr Thr Arg Gly Pro His 965 970 975

Lys Asp Glu Phe Cys Pro Ser Thr Asp Thr Ser Phe Asn Cys Leu Lys 980 985 990

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Ile	Phe 1115	Thr	Leu	Ala	Leu	Cys 1120	Phe	Val	Pro	Thr	Phe 1125	Val	Val	Cys
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Lys Leu Ala Thr Glu Ala Ala Met Val Ser Glu Pro Ser Cys Pro 1280 1285 1290

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gcc ttc t Ala Phe T 35	_	_		_			-			-		203
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tac acc g Tyr Thr G	_		_	-	_		_	_	_		_	347
tca ctg g Ser Leu G 1		_	_	e Thr	_							395
cca gcc t Pro Ala C 115												443
agc ccc a Ser Pro A 130	_	_					-		_			491
ggg gct g Gly Ala G					_			_			_	539
cat agc t His Ser P	_		_	_		_	_	_			_	587
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	atc Ile 355	_				_		_			_	_	-			1163
	gtc Val															1211
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	690			695				700			705	
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					Gly 999							2267
					ccc Pro 745							2315
					gly ggg							2363
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Ŧ					gtc Val							2507
					gag Glu 825							2555
					ctg Leu							2603
					ttc Phe							2651
					cag Gln							2699
					ttt Phe							2747
	-	_		_	acc Thr 905	_				_		2795
					tcc Ser	-	_					2843
					gcc Ala							2891

ctg gcc atc cct gcc tcc tcc tgg gtg gat gac ttc att gac tgg ctg Leu Ala Ile Pro Ala Ser Ser Trp Val Asp Asp Phe Ile Asp Trp Leu 950 955 960	2939
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ccc aaa ggc ggc ctg gca gca tac agc acc tct gtg aac ttg act Pro Lys Gly Gly Leu Ala Ala Tyr Ser Thr Ser Val Asn Leu Thr 1025 1030 1035	3173
tca gat ggc cag gtt tta gac aca gtt gcc att ctg tca ccc agg Ser Asp Gly Gln Val Leu Asp Thr Val Ala Ile Leu Ser Pro Arg 1040 1045 1050	3218
ctg gag tac agt ggc aca atc tcg gct cac tgc aac ctc tac ctc Leu Glu Tyr Ser Gly Thr Ile Ser Ala His Cys Asn Leu Tyr Leu 1055 1060 1065	3263
ctg gat tca gcc tcc agg ttc atg gcc tat cac aag ccc ctg aaa Leu Asp Ser Ala Ser Arg Phe Met Ala Tyr His Lys Pro Leu Lys 1070 1075 1080	3308
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gca gcc aac atc act gct gac ctg cgg aaa gtg cct gga aca gac Ala Ala Asn Ile Thr Ala Asp Leu Arg Lys Val Pro Gly Thr Asp 1100 1105 1110	3398
ccg gct ttt gag gtc ttc ccc tac acg atc acc aat gtg ttt tat Pro Ala Phe Glu Val Phe Pro Tyr Thr Ile Thr Asn Val Phe Tyr 1115 1120 1125	3443
gag cag tac ctg acc atc ctc cct gag ggg ctc ttc atg ctc agc Glu Gln Tyr Leu Thr Ile Leu Pro Glu Gly Leu Phe Met Leu Ser 1130 1135 1140	3488
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tac Tyr 1190										tcg Ser 1200					3668
tct Ser 1205										tcc Ser 1215					3713
acc Thr 1220										gag Glu 1230					3758
atg Met 1235		_		_		_			-	atg Met 1245			_	cct Pro	3803
ggc Gly 1250										cag Gln 1260					3848
ttc Phe 1265										ctg Leu 1275	_		_	_	3893
cat His 1280		_			_					agc Ser 1290		gtg Val			3938
gac Asp 1295										aag Lys 1305					3983
gcg Ala 1310		_	_	_	_		_		_	cca Pro 1320					4028
cga Arg 1325	Val									aac Asn 1335					4073
ggt Gly 1340	Ser				_	Gly	_		_	aac Asn 1350		_			4118
aat Asn 1355	Gly			ttc Phe	tga	tacag	gcca	ga g	gecet	gtet	agg	ctcta	atg		4166
gccc	tgaa	сс а	aagg	gttai	t ggg	gatci	ttc (	cttg	tgact	g cc	cctt	gaca	cac	gecetee	4226
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Cys Ala Phe Tyr Asp Glu Cys Gly Lys Asn Pro Glu Leu Ser Gly Ser 35 40 45

Leu Met Thr Leu Ser Asn Val Ser Cys Leu Ser Asn Thr Pro Ala Arg 50 55 60

Lys Ile Thr Gly Asp His Leu Ile Leu Leu Gln Lys Ile Cys Pro Arg 65 70 75 80

Leu Tyr Thr Gly Pro Asn Thr Gln Ala Cys Cys Ser Ala Lys Gln Leu 85 90 95

Val Ser Leu Glu Ala Ser Leu Ser Ile Thr Lys Ala Leu Leu Thr Arg

100	105	110

Cys	Pro	Ala 115	Cys	Ser	Asp	Asn	Phe 120	Val	Asn	Leu	His	Cys 125	His	Asn	Thr
Cys	Ser 130	Pro	Asn	Gln	Ser	Leu 135	Phe	Ile	Asn	Val	Thr 140	Arg	Val	Ala	Gln
Leu 145	Gly	Ala	Gly	Gln	Leu 150	Pro	Ala	Val	Val	Ala 155	Tyr	Glu	Ala	Phe	Tyr 160
Gln	His	Ser	Phe	Ala 165	Glu	Gln	Ser	Tyr	Asp 170	Ser	Cys	Ser	Arg	Val 175	Arg
Val	Pro	Ala	Ala 180	Ala	Thr	Leu	Ala	Val 185	Gly	Thr	Met	Cys	Gly 190	Val	Tyr
Gly	Ser	Ala 195	Leu	Cys	Asn	Ala	Gln 200	Arg	Trp	Leu	Asn	Phe 205	Gln	Gly	Asp
Thr	Gly 210	Asn	Gly	Leu	Ala	Pro 215	Leu	Asp	Ile	Thr	Phe 220	His	Leu	Leu	Glu
Pro 225	Gly	Gln	Ala	Val	Gly 230	Ser	Gly	Ile	Gln	Pro 235	Leu	Asn	Glu	Gly	Val 240
Ala	Arg	Cys	Asn	Glu 245	Ser	Gln	Gly	Asp	Asp 250	Val	Ala	Thr	Cys	Ser 255	Cys
Gln	Asp	Cys	Ala 260	Ala	Ser	Cys	Pro	Ala 265	Ile	Ala	Arg	Pro	Gln 270	Ala	Leu
Asp	Ser	Thr 275	Phe	Tyr	Leu	Gly	Gln 280	Met	Pro	Gly	Ser	Leu 285	Val	Leu	Ile
Ile	Ile 290	Leu	Cys	Ser	Val	Phe 295	Ala	Val	Val	Thr	Ile 300	Leu	Leu	Val	Gly
Phe 305	Arg	Val	Ala	Pro	Ala 310	Arg	Asp	Lys	Ser	Lys 315	Met	Val	Asp	Pro	Lys 320
Lys	Gly	Thr	Ser	Leu 325	Ser	Asp	Lys	Leu	Ser 330	Phe	Ser	Thr	His	Thr 335	Leu
Leu	Gly	Gln	Phe 340	Phe	Gln	Gly	Trp	Gly 345	Thr	Trp	Val	Ala	Ser 350	Trp	Pro

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Glu	Glu	Ala 595	Phe	Leu	Glu	Glu	Met 600	Arg	Ala	Phe	Gln	Arg 605	Arg	Met	Ala
Gly	Met 610	Phe	Gln	Val	Thr	Phe 615	Met	Ala	Glu	Arg	Ser 620	Leu	Glu	Asp	Glu
Ile 625	Asn	Arg	Thr	Thr	Ala 630	Glu	Asp	Leu	Pro	Ile 635	Phe	Ala	Thr	Ser	Tyr 640
Ile	Val	Ile	Phe	Leu 645	Tyr	Ile	Ser	Leu	Ala 650	Leu	Gly	Ser	Tyr	Ser 655	Ser
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Tyr	Leu 690	Gly	Ile	Arg	Ser	Ser 695	Leu	Val	Ile	Leu	Gln 700	Val	Val	Pro	Phe
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Tyr	Gln	Arg	Leu	Pro 725		Arg	Pro	Gly	Glu 730	Pro	Arg	Glu	Val	His 735	Ile
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Ser	Glu	Ala 755		Cys	Phe	Phe	Leu 760		Ala	Leu	Thr	Pro 765		Pro	Ala
Val	Arg 770		Phe	: Ala	Leu	Thr 775		Gly	Leu	Ala	Val 780		Leu	Asp	Phe
Leu 785		ı Glr	n Met	: Ser	790		Val	Ala	Leu	Leu 795		Leu	Asp	Ser	Lys 800
Arg	g Glr	ı Glu	ı Ala	Ser 805		Leu	ı Asp	Val	Cys 810		Cys	Val	Lys	Pro 815	Gln
Glı	ı Leı	ı Pro	Pro 820		Gly	g Glr	ı Gly	Glu 825		Leu	Leu	Leu	Gly 830		Phe

- Gln Lys Ala Tyr Ala Pro Phe Leu Leu His Trp Ile Thr Arg Gly Val 835 840 845
- Val Leu Leu Phe Leu Ala Leu Phe Gly Val Ser Leu Tyr Ser Met 850 855 860
- Cys His Ile Ser Val Gly Leu Asp Gln Glu Leu Ala Leu Pro Lys Asp 865 870 875 880
- Ser Tyr Leu Leu Asp Tyr Phe Leu Phe Leu Asn Arg Tyr Phe Glu Val 885 890 895
- Gly Ala Pro Val Tyr Phe Val Thr Thr Leu Gly Tyr Asn Phe Ser Ser 900 905 910
- Glu Ala Gly Met Asn Ala Ile Cys Ser Ser Ala Gly Cys Asn Asn Phe 915 920 925
- Ser Phe Thr Gln Lys Ile Gln Tyr Ala Thr Glu Phe Pro Glu Gln Ser 930 935 940
- Tyr Leu Ala Ile Pro Ala Ser Ser Trp Val Asp Asp Phe Ile Asp Trp 945 950 955 960
- Leu Thr Pro Ser Ser Cys Cys Arg Leu Tyr Ile Ser Gly Pro Asn Lys 965 970 975
- Asp Lys Phe Cys Pro Ser Thr Val Asn Ser Leu Asn Cys Leu Lys Asn 980 985 990
- Cys Met Ser Ile Thr Met Gly Ser Val Arg Pro Ser Val Glu Gln Phe 995 1000 1005
- His Lys Tyr Leu Pro Trp Phe Leu Asn Asp Arg Pro Asn Ile Lys 1010 1015 1020
- Cys Pro Lys Gly Gly Leu Ala Ala Tyr Ser Thr Ser Val Asn Leu 1025 1030 1035
- Thr Ser Asp Gly Gln Val Leu Asp Thr Val Ala Ile Leu Ser Pro 1040 1045 1050
- Arg Leu Glu Tyr Ser Gly Thr Ile Ser Ala His Cys Asn Leu Tyr 1055 1060 1065
- Leu Leu Asp Ser Ala Ser Arg Phe Met Ala Tyr His Lys Pro Leu

1070	1075	1080

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Asp	Pro 1115	Ala	Phe	Glu	Val	Phe 1120		Tyr	Thr	Ile	Thr 1125	Asn	Val	Phe
Tyr	Glu 1130	Gln	Tyr	Leu	Thr	Ile 1135	Leu	Pro	Glu	Gly	Leu 1140		Met	Leu
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Gly	Leu 1160	Asp	Leu	Arg	Ser	Gly 1165	Leu	Leu	Asn	Leu	Leu 1170	Ser	Ile	Val
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Met	Ser - 1205	Val	Glu	Phe	Val	Ser 1210	His	Ile	Thr	Arg	Ser 1215	Phe	Ala	Ile
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Ser	Met 1235	Gly	Ser	Ala	Val	Phe 1240	Ala	Gly	Val	Ala	Met 1245	Thr	Asn	Leu
Pro	Gly 1250	Ile	Leu	Val	Leu	Gly 1255	Leu	Ala	Lys	Ala	Gln 1260	Leu	Ile	Gln
Ile	Phe 1265	Phe	Phe	Arg	Leu	Asn 1270	Leu	Leu	Ile	Thr	Leu 1275	Leu	Gly	Leu
Leu	His 1280	Gly	Leu	Val	Phe	Leu 1285	Pro	Val	Ile	Leu	Ser 1290	Tyr	Val	Gly
Pro	Asp 1295	Val	Asn	Pro	Ala	Leu 1300	Ala	Leu	Glu	Gln	Lys 1305	Arg	Ala	Glu

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Ser Arg Val Ser Thr Ala Asp Asn Ile Tyr Val Asn His Ser Phe 1325 1330 1335

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Asn Asn Gly Arg Gln Phe 1355

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